C57BL6			C3H/HeN			
	Fold change (chronic v	Functional Annotation		Fold change (chronic v	Functional Annotation	
Gene name	resolved)	5 (Gene name	resolved)		
Ubd	119.405	Proteosome	BC100530	144.401	Unknown	
Saa3	112.203	Inflammatory Response	S100a8	88.7327	Cytokine/lon Binding	
S100a8	85.4193	Cytokine/ lon Binding	Mmp7	87.0008	Ion Binding	
Mmp7	77.6183	Ion Binding	Ubd	77.6292	Proteosome	
RegIIIγ	66.2146	Inflammatory Response	S100a9	74.4965	Cytokine/ Ion Binding	
S100a9	44.9876	Cytokine/ Ion Binding	RegIIIγ	62.1077	Inflammatory Response	
lgk-V19-14	31.1745	Immunoglobulin	Cxcl2	39.3707	Cytokine	
lgk-V28	27.7003	Immunoglobulin	II1b	38.9361	Cytokine	
Fam3b	25.7301	Cytokine	Sprr2f	38.7662	Tissue Remodeling	
Plekhs1	24.3332	Tissue Remodeling	Saa3	36.8235	Inflammatory Response	
Duox2	23.492	Ion Binding	Cxcl5	34.2451	Cytokine	
Gm5571	22.2598	Immunoglobulin	ll1a	33.0689	Cytokine	
Sprr2f	22.0386	Tissue Remodeling	Gabrp	31.6717	Ion Binding	
AA467197	21.9697		Sprr2d	31.4934	Tissue Remodeling	
Cxcl9	21.46	Cytokine	Fam3b	29.3965	Cytokine	
Cxcl2	17.698	Cytokine	Cxcr2	27.6748	Cytokine	
Pigr	17.1008	Immunoglobulin	Tmprss11g	26.263	Tissue Remodeling	
Tmprss11g	15.7865	Tissue Remodeling	Lcn2	25.9349	Ion Binding	
Nos2	15.644	Inflammatory Response	II1f9	24.0818	Cytokine	
Il1a	15.0219	Cytokine	Sprr2g	23.4676	Tissue Remodeling	